	Application No.	Applicant(s)	0
	09/407,605	MILLER ET AL.	
Notice of Allowability	Examiner	Art Unit	
	Chih-Min Kam	1653	
The MAILING DATE of this communication appear All claims being allowable, PROSECUTION ON THE MERITS IS herewith (or previously mailed), a Notice of Allowance (PTOL-85) NOTICE OF ALLOWABILITY IS NOT A GRANT OF PATENT RI	ears on the cover sheet wi (OR REMAINS) CLOSED in or other appropriate comm IGHTS. This application is:	th the correspondence address n this application. If not included unication will be mailed in due course. THIS	
1. This communication is responsive to 8/9/04.			
2. The allowed claim(s) is/are <u>81-94,96-108 and 110-112</u> .			
3. The drawings filed on are accepted by the Examine	r.		
 4. ☐ Acknowledgment is made of a claim for foreign priority una a) ☐ All b) ☐ Some* c) ☐ None of the: 1. ☐ Certified copies of the priority documents have 2. ☐ Certified copies of the priority documents have 3. ☐ Copies of the certified copies of the priority documents have International Bureau (PCT Rule 17.2(a)). * Certified copies not received: 	been received. been received in Application	on No	
Applicant has THREE MONTHS FROM THE "MAILING DATE" on noted below. Failure to timely comply will result in ABANDONM THIS THREE-MONTH PERIOD IS NOT EXTENDABLE.	of this communication to file IENT of this application.	a reply complying with the requirements	
5. A SUBSTITUTE OATH OR DECLARATION must be submit INFORMAL PATENT APPLICATION (PTO-152) which give	itted. Note the attached EXA es reason(s) why the oath o	MINER'S AMENDMENT or NOTICE OF declaration is deficient.	
 CORRECTED DRAWINGS (as "replacement sheets") mus (a) including changes required by the Notice of Draftspers 1) hereto or 2) to Paper No./Mail Date 3/29/05 (b) including changes required by the attached Examiner's Paper No./Mail Date 20041025. Identifying indicia such as the application number (see 37 CFR 1. each sheet. Replacement sheet(s) should be labeled as such in the 	on's Patent Drawing Review 1. 5 Amendment / Comment or 84(c)) should be written on the header according to 37 CF	in the Office action of ne drawings in the front (not the back) of R 1.121(d).	
 DEPOSIT OF and/or INFORMATION about the deposit attached Examiner's comment regarding REQUIREMENT F 	sit of BIOLOGICAL MATE FOR THE DEPOSIT OF BIO	RIAL must be submitted. Note the PLOGICAL MATERIAL.	
Attachment(s) 1. ☐ Notice of References Cited (PTO-892) 2. ☐ Notice of Draftperson's Patent Drawing Review (PTO-948) 3. ☐ Information Disclosure Statements (PTO-1449 or PTO/SB/08 Paper No./Mail Date 4. ☐ Examiner's Comment Regarding Requirement for Deposit of Biological Material	6. ⊠ Interview Sı Paper No./ 8), 7. ⊠ Examiner's	formal Patent Application (PTO-152) Jammary (PTO-413), Mail Date <u>20041025</u> . Amendment/Comment Statement of Reasons for Allowance	

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An **Examiner's Amendment** to the record appears below. Should the changes and/or additions be unacceptable to applicant, an amendment may be filed as provided by 37 CFR 1.312. To ensure consideration of such an amendment, it MUST be submitted no later than the payment of the issue fee.

Authorization for this examiner's amendment was given in a telephone interview with Todd Garcia on October 25, 2004.

Examiner's Amendments to the Claims:

Claims 81, 85, 89, 97, 100 and 103 have been amended as follows:

- 81. (Currently amended) A synthetic nucleic acid sequence which encodes <u>a</u> human Factor VIII or [a functional portion thereof], wherein at least one non-common codon or less-common codon has been replaced by a common codon encoding the same amino acid residue as the non-common or less-common codon and wherein the synthetic nucleic acid has a continuous stretch of at least 150 codons all of which are common codons, <u>and</u> wherein by a common codon is meant the most common codon encoding each particular amino acid residue in highly expressed human genes as shown in Figures 14A[-] <u>and 14B</u>.
- 85. (Currently amended) A synthetic nucleic acid sequence which encodes <u>a</u> human Factor VIII [or a functional portion thereof], wherein at least one non-common codon or less-common codon has been replaced by a common codon encoding the same amino acid residue as the non-common or less-common codon and wherein the synthetic nucleic acid has a continuous stretch of common codons which comprise at least 60% of the codons of the synthetic nucleic acid sequence, <u>and</u> wherein by a common codon is meant the most common codon encoding each particular amino acid residue in highly expressed human genes as shown in Figures 14A[-] and 14B.
- 89. (Currently amended) A synthetic nucleic acid sequence which encodes <u>a</u> human Factor VIII [or a functional portion thereof], wherein at least one non-common codon or less-common codon has been replaced by a common codon encoding the same amino acid residue as the non-common or less-common codon and wherein at least 98% or more of the codons in the sequence encoding the Factor VIII are common codons and the Factor VIII is at least 90 amino

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acid residues in length, and wherein by a common codon is meant the most common codon encoding each particular amino acid residue in highly expressed human genes as shown in Figures 14A[-] and 14B.

- 97. (Currently amended) A synthetic nucleic acid sequence which encodes human Factor IX, wherein at least one non-common codon or less-common codon has been replaced by a common codon encoding the same amino acid residue as the non-common or less-common codon and wherein the synthetic nucleic acid has a continuous stretch of at least 150 codons all of which are common codons, and wherein by a common codon is meant the most common codon encoding each particular amino acid residue in highly expressed human genes as shown in Figures 14A[-] and 14B.
- 100. (Currently amended) A synthetic nucleic acid sequence which encodes human Factor IX, wherein at least one non-common codon or less-common codon has been replaced by a common codon encoding the same amino acid residue as the non-common or less-common codon and wherein the synthetic nucleic acid has a continuous stretch of common codons which comprise at least 60% of the codons of the synthetic nucleic acid sequence, and wherein by a common codon is meant the most common codon encoding each particular amino acid residue in highly expressed human genes as shown in Figures 14A[-] and 14B.
- 103. (Currently amended) A synthetic nucleic acid sequence which encodes human Factor IX, wherein at least one non-common codon or less-common codon has been replaced by a common codon encoding the same amino acid residue as the non-common or less-common codon and wherein at least 98% or more of the codons in the sequence encoding the Factor IX are common codons and the Factor IX is at least 90 amino acid residues in length, and wherein by a common codon is meant the most common codon encoding each particular amino acid residue in highly expressed human genes as shown in Figures 14A[-] and 14B.

Objection to Drawings

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Figures 14A and 14B should be designated by a legend such as --Prior Art-- because only that which is old is illustrated. See MPEP § 608.02(g). Corrected drawings in compliance with 37 CFR 1.121(d) are required in reply to the Office action to avoid abandonment of the application. The replacement sheet(s) should be labeled "Replacement Sheet" in the page header (as per 37 CFR 1.121(d)) so as not to obstruct any portion of the drawing figures. The objection to the drawings will not be held in abeyance.

The following is an **Examiner's Statement of Reasons for Allowance**: The following reference appears to be the closest art to the claimed invention. Seed *et al.* teach a synthetic gene encoding a protein such as factor VIII normally expressed in a mammalian cell or other eukaryotic cells, wherein at least one non-preferred or less preferred codon in the normal gene encoding the protein has been replaced by a preferred codon encoding the same amino acid, where the preferred codon is indicated in the highly expressed human genes; and at least 10%, 20%, 30%, 40%, 50%, 60%, 70%, 80% or 90% of the non-preferred codons in the natural gene are replaced with preferred codons. However, Seed *et al.* do not disclose the synthetic nucleic acid having a continuous stretch of at least 150 common codons or of 60% of the codons of the synthetic nucleic acid sequence, or at least 98% of more of the codons in the sequence encoding the protein (e.g., factor VIII or factor IX) are common codons, and the protein is at least 90 amino acid residues in length. Therefore, the claims are allowable over the art of record.

Any comments considered necessary by applicant must be submitted no later than the payment of the issue fee and, to avoid processing delays, should preferably accompany the issue fee. Such submissions should be clearly labeled "Comments on Statement of Reasons for Allowance."

Any inquiry concerning this communication or earlier communications from the examiner should be directed to Chih-Min Kam whose telephone number is (571) 272-0948. The examiner can normally be reached on 8.00-4:30, Mon-Fri.

If attempts to reach the examiner by telephone are unsuccessful, the examiner's supervisor, Jon Weber can be reached at 571-272-0925. The fax phone number for the organization where this application or proceeding is assigned is 703-872-9306.

Information regarding the status of an application may be obtained from the Patent Application Information Retrieval (PAIR) system. Status information for published applications may be obtained from either Private PAIR or Public PAIR. Status information for unpublished applications is available through Private PAIR only. For more information about the PAIR system, see http://pair-direct.uspto.gov. Should you have questions on access to the Private PAIR system, contact the Electronic Business Center (EBC) at 866-217-9197 (toll-free).

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Chih-Min Kam, Ph. D. CITL
Patent Examiner

CMK

October 25, 2004

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